

FIGURE 1A

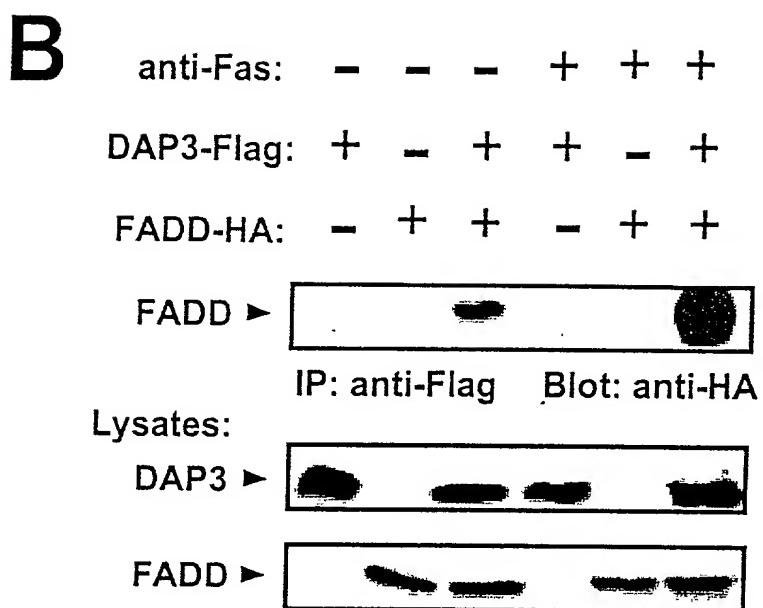


FIGURE 1B

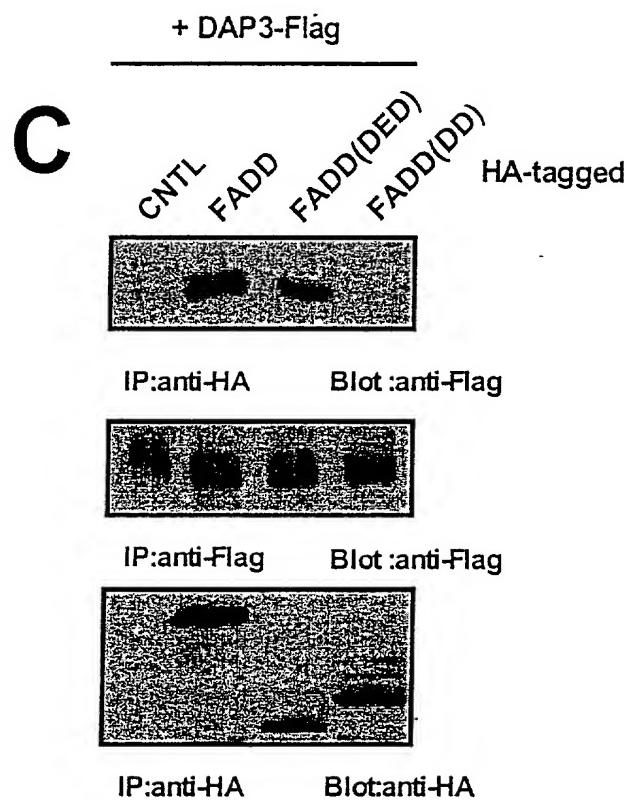


FIGURE 1C

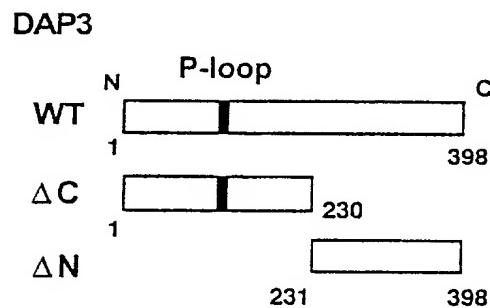
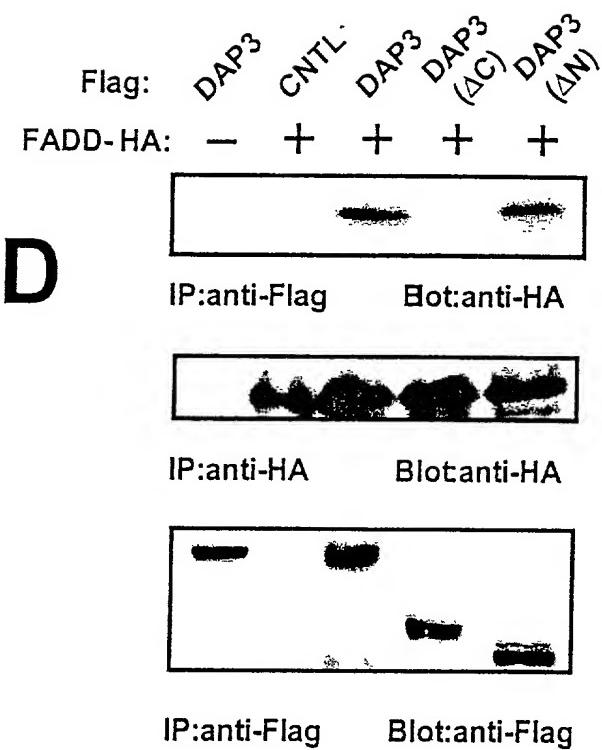


FIGURE 1D

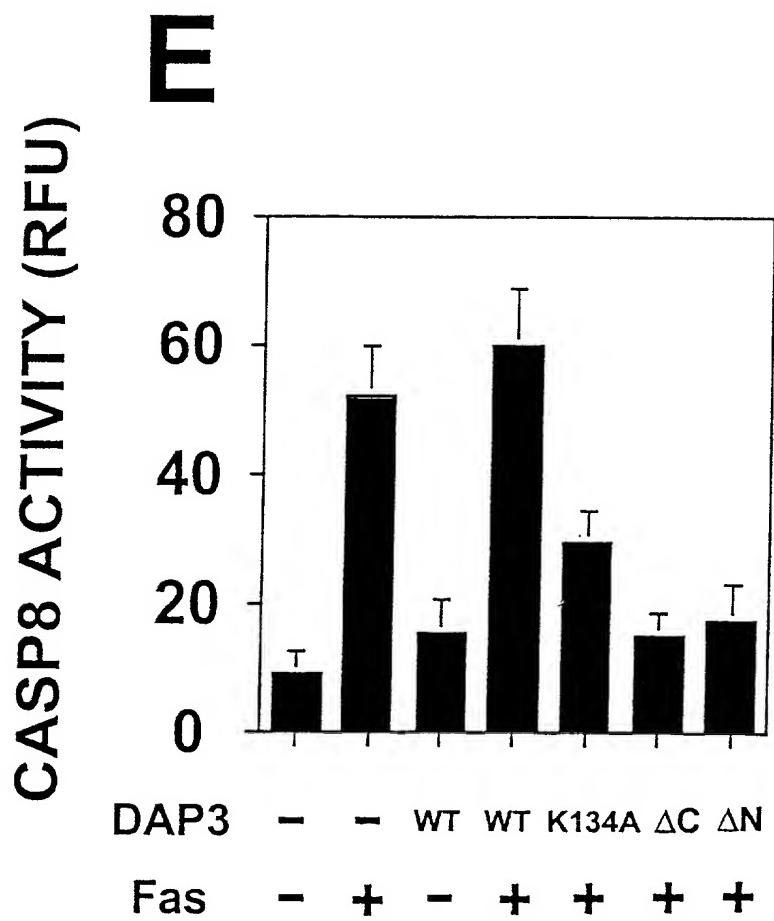


FIGURE 1E

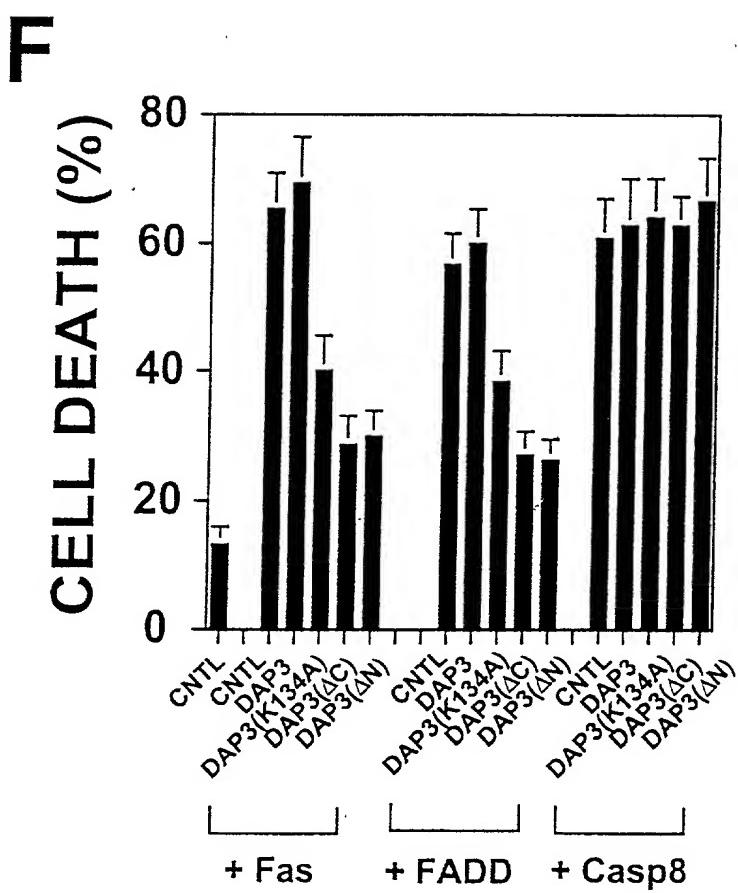


FIGURE 1F

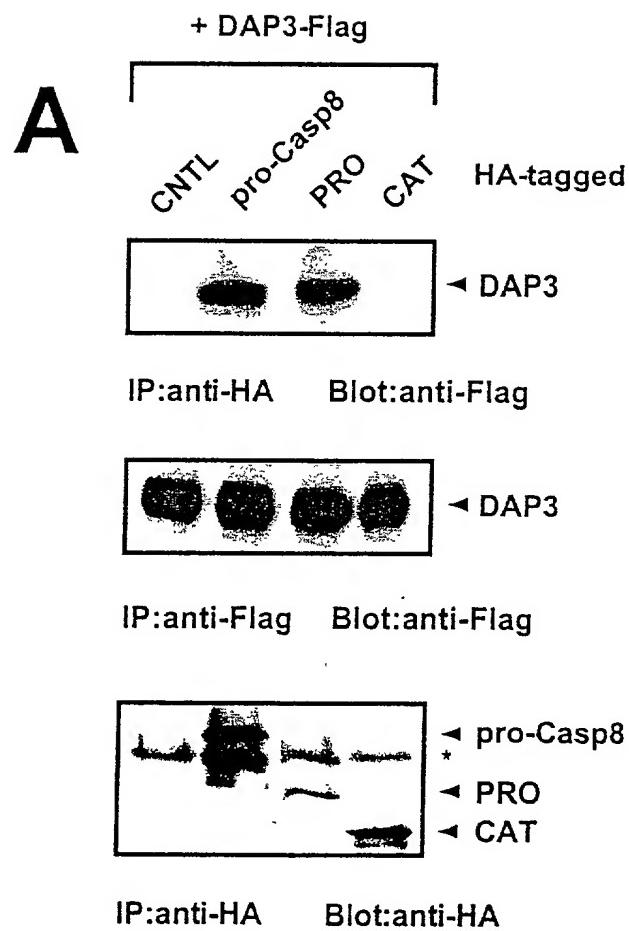


FIGURE 2A

B

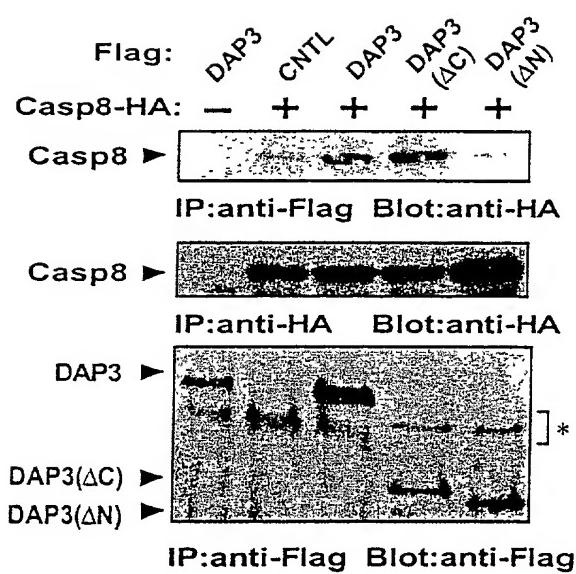


FIGURE 2B

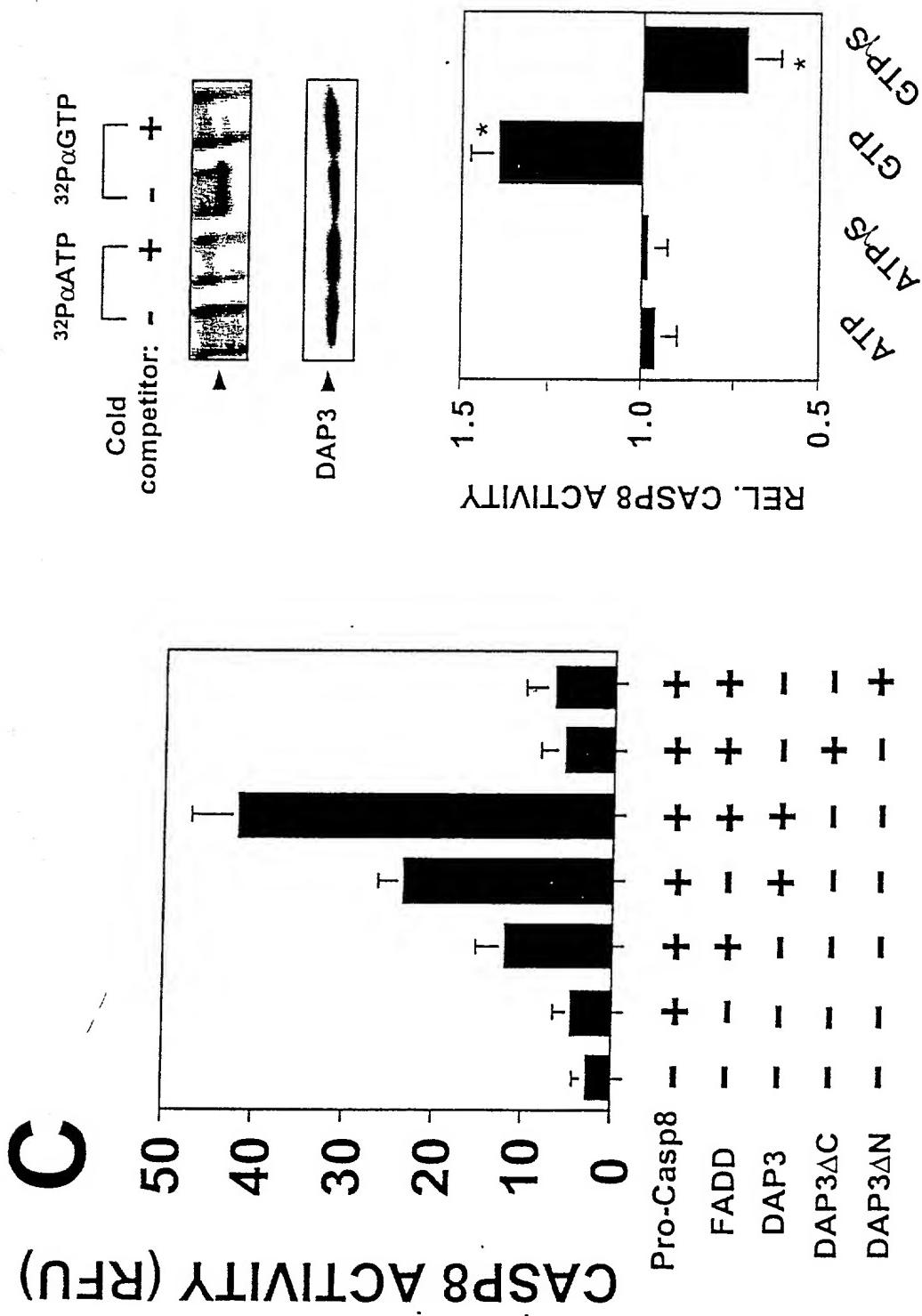


FIGURE 2C

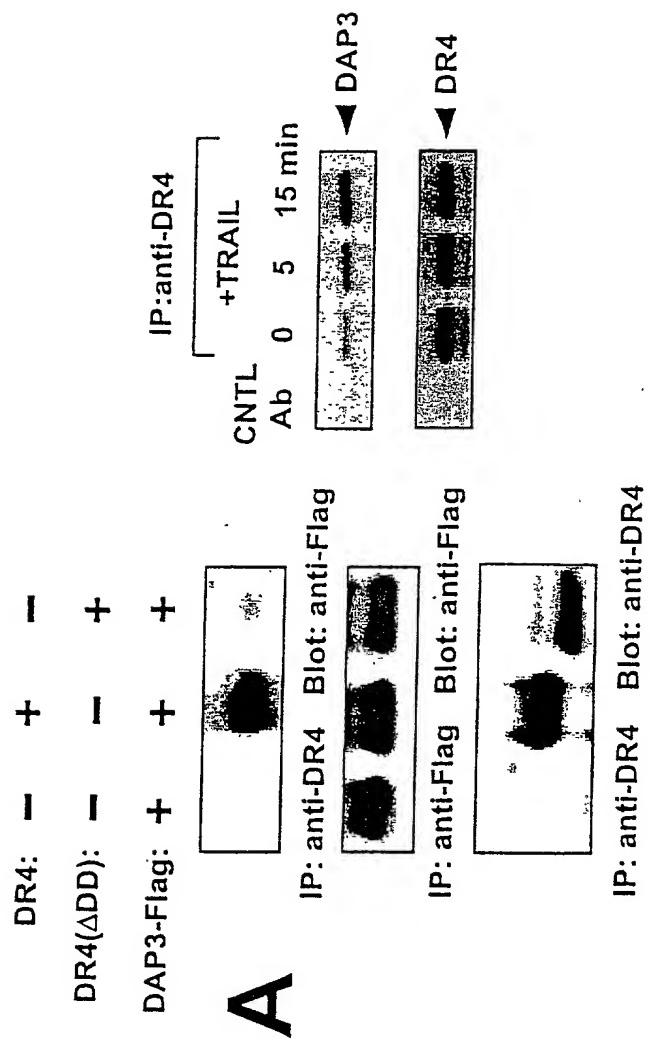


FIGURE 3A

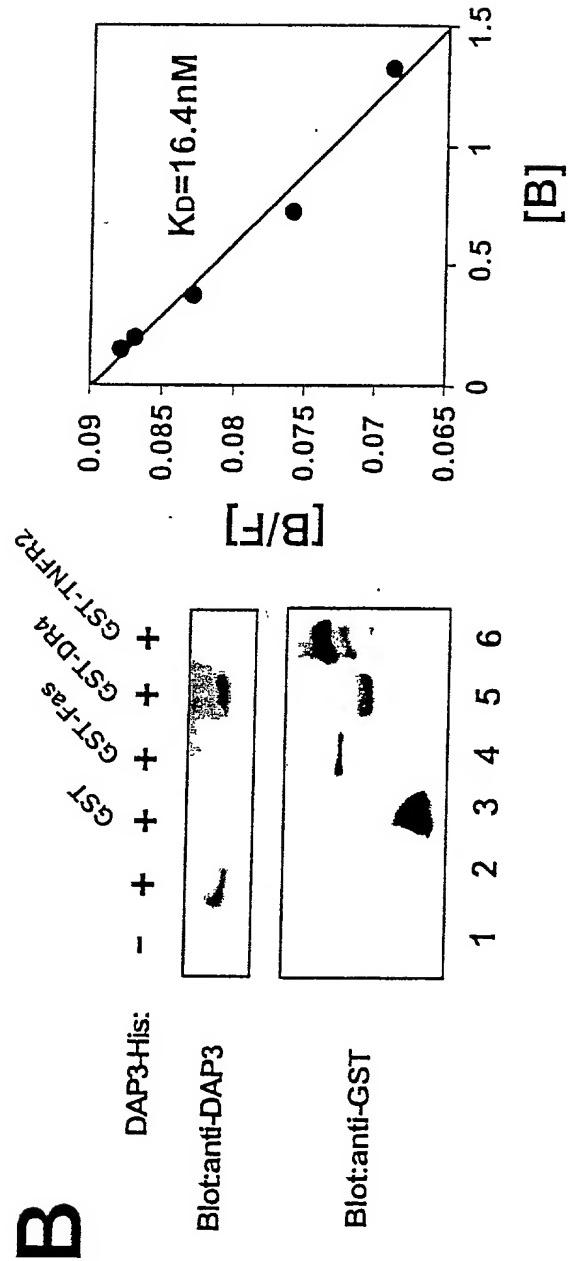


FIGURE 3B

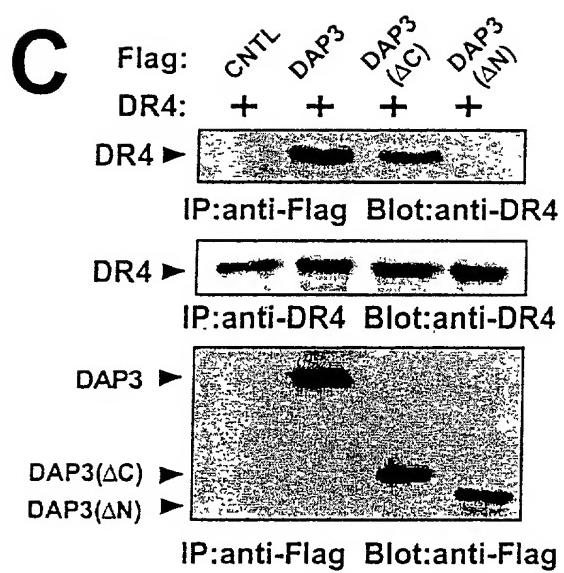


FIGURE 3C

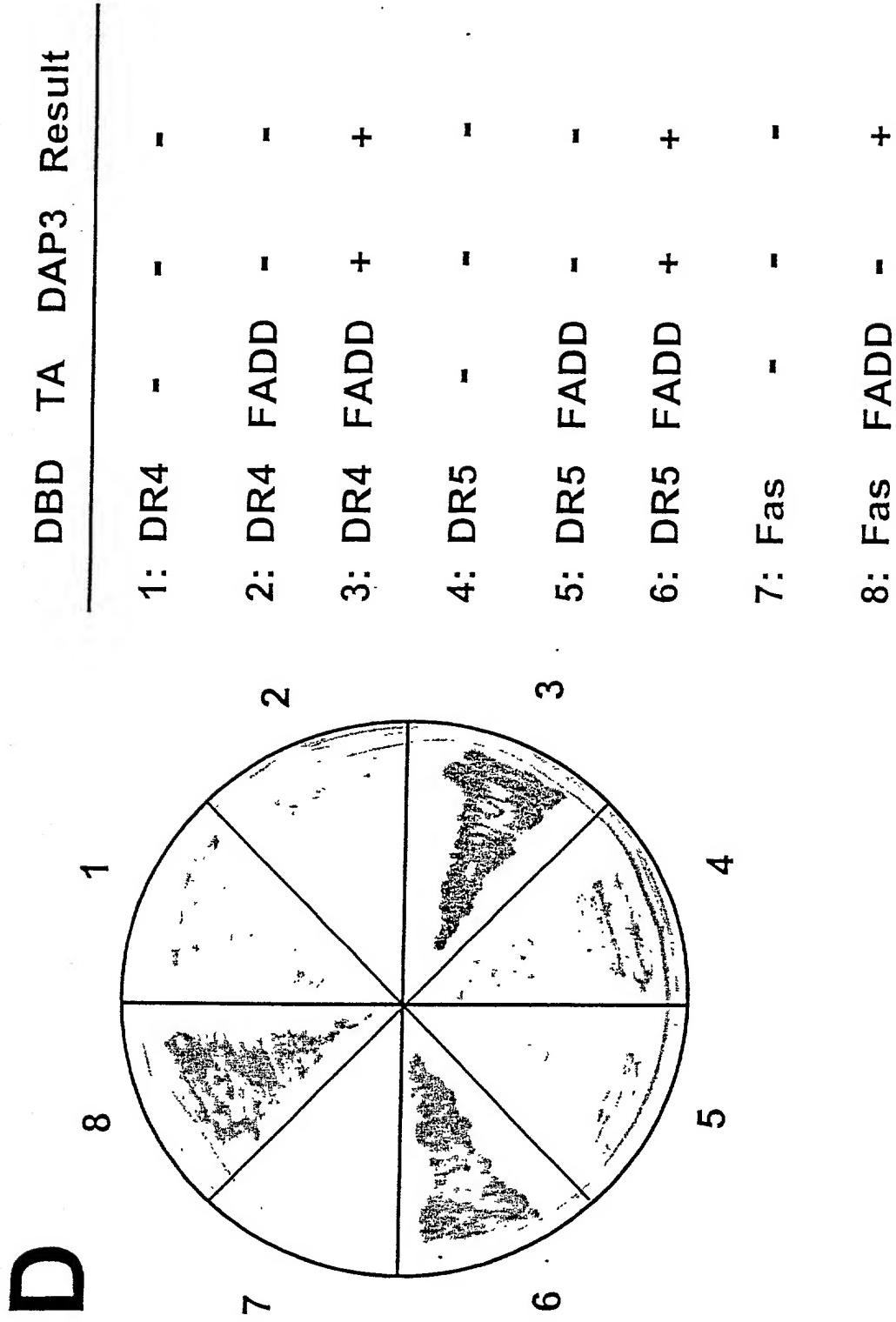


FIGURE 3D

E

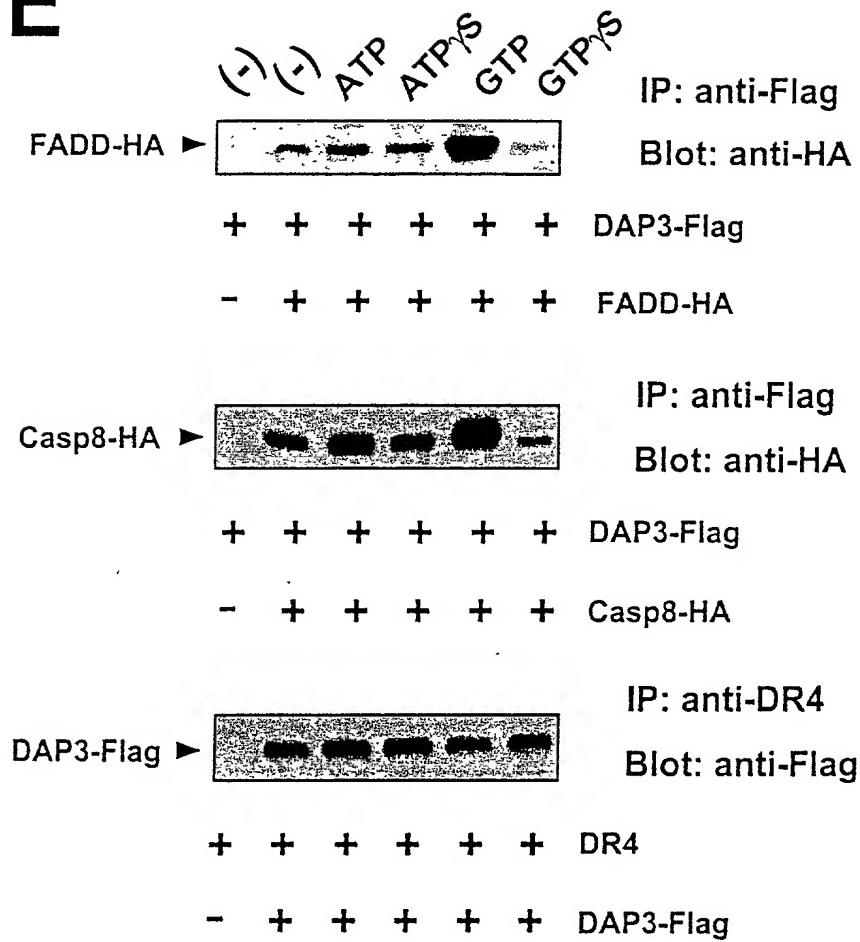


FIGURE 3E

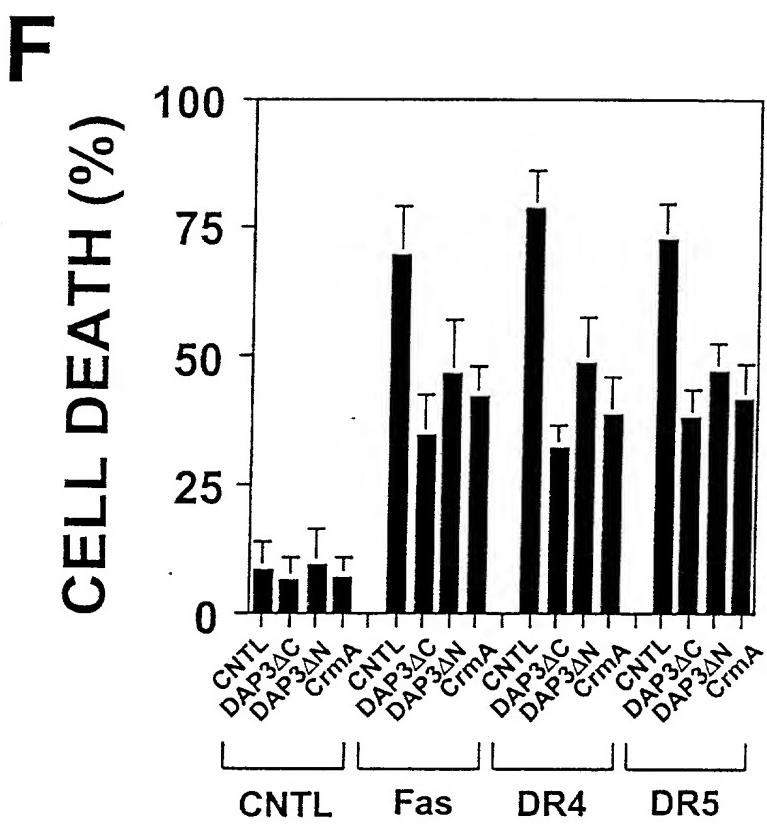


FIGURE 3F

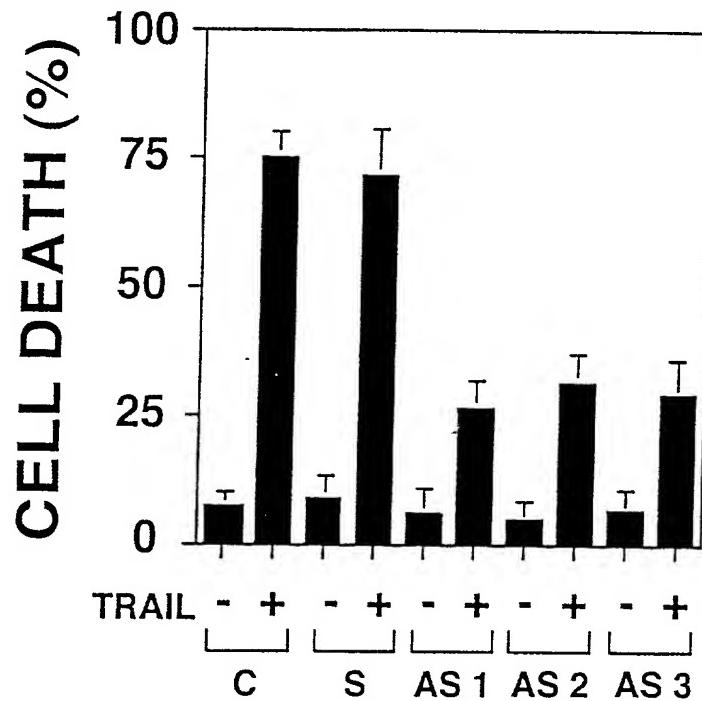
G

DAP3
agtccctcga ctcacgtgca aggatgtgc tgaaaggaata

antisense 1

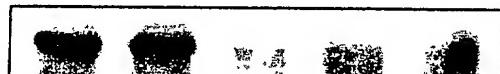
antisense 2

antisense 3



C S AS 1 AS 2 AS 3

DAP3 ▶



DR4 ▶

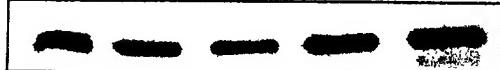


FIGURE 3G

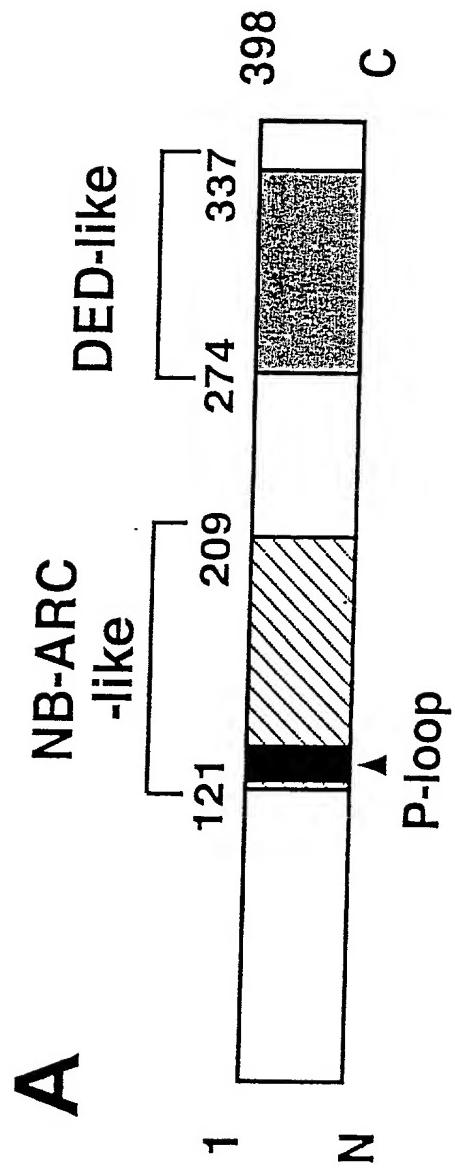


FIGURE 4A

B

| | | | | | | |
|-----|---|---------------------------------------|-------------------------------------|-----------------------------------|-------------|--------|
| 115 | T S F A Y P A I R Y L | L Y G E K G K I T S S L C H V | - - - - - | F C A K Q D W L | - - - | DAP3 |
| 142 | K N L K G - E P G W V T | H G M A G C G K S V A S G A | - - - - - | P G G Y V W V S V G K | Q | Apa1IN |
| 146 | C D L D S F | H G R A G S G K S V A S G A | - - - - - | V W I K D S G T | CED4 | |
| 152 | - - - - - | - - - - - | - - - - - | - - - - - | - - - | - |
| 191 | D K S G L L M K L Q N C T R L | D Q D I E S F S Q R L P L N E | - - - - - | H P D A H L W V K N C R D E | E R Q S S Y | DAP3 |
| 196 | A P K S T F D L F D I L L M L K S | D L - L N F P S M E H V | - - - - - | E Y K D R L R K H P R | W V D R P | Apa1IN |
| 174 | N K Q R F D Q P | I E A S T - - - - - | W - L K N F K - - - - - | I T N E R F V N Q I K V Q E K Y V | W N K R | DAP3 |
| 238 | S L D D V W D S - - - - - | W V L K N F D S Q C A I E | I T T R D X S V M G P K Y V | V P V E | Apa1IN | |
| 245 | F D D V W Q E E T I R W A - Q E L R L R C - - - - - | E V T T R D V E V S N A A S Q T C E F | I T T R D V E V S N A A S Q T C E F | V E | CED4 | |

C

| | | | | | | |
|-----|---|---|-----------------|-------------------------|-------|------------|
| 268 | W W G R T F L K R E D | S P L A E E E L A L W H N L | - - - - - | R K M W K N D W H G G V | S A L | DAP3 DED |
| 1 | M D F S R N L Y D I S E | E S E D L A S L K F L S L D | - - - - - | P I K D A L M L F Q R L | Q | Casp8 DED1 |
| 98 | H L L I R V W L Y D I S E | R S F K F L L Q E E S K C K L D | - - - - - | D D M L L D F I E M E | E | Casp8 DED2 |
| 17 | K V S F R E K L L I E S N L G V | D V A V N L K F L C I G L V P N K K L E K S S S D A F H | - - - - - | Casp10 DED1 | | |
| 121 | V S S L F R N L L Y E S E G D S E G D S E G D S S S S E L | N L K D S I P K T E M T S - - - - - | L S F L A F L E | Casp10 DED2 | | |
| 1 | M D P F L V L L H S V S S S S S S S E L | I E L K F L C L G R V G K J F K L E R V Q S G L D L F S M L L | - - - - - | F ADD DED | | |
| 312 | Q T G S L F K P R K A X L P Q E L G K E G F D A L | - - - - - | - - - - - | - - - - - | - | - |
| 50 | E K R M L E E S N L S F K E L L F R I N R L D L L | - - - - - | - - - - - | - - - - - | - | - |
| 148 | K R V I L G E G K L D I L K R V A C A Q I N K - S L L | - - - - - | - - - - - | - - - - - | - | - |
| 67 | A S E D L S E E D L P F L A E L L Y I I R Q K K L L | - - - - - | - - - - - | - - - - - | - | - |
| 167 | K Q G K D E D N L T C E D L C K T V P - K L L R | - - - - - | - - - - - | - - - - - | - | - |
| 51 | E Q N D L E P G H T E L L R E L L A S R R H D L L R | - - - - - | - - - - - | - - - - - | - | - |

FIGURE 4B, C

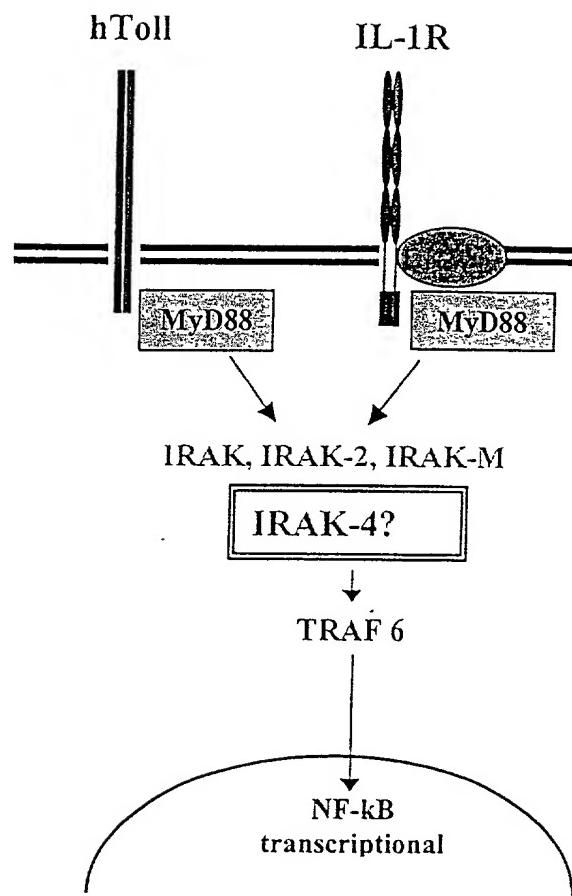


FIGURE 5

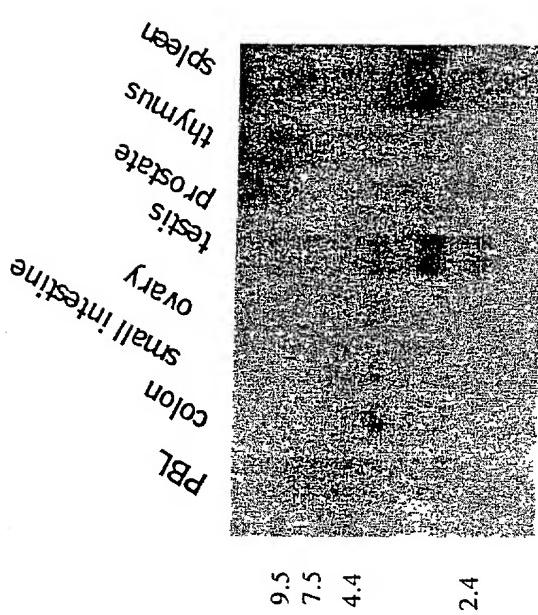


FIGURE 6

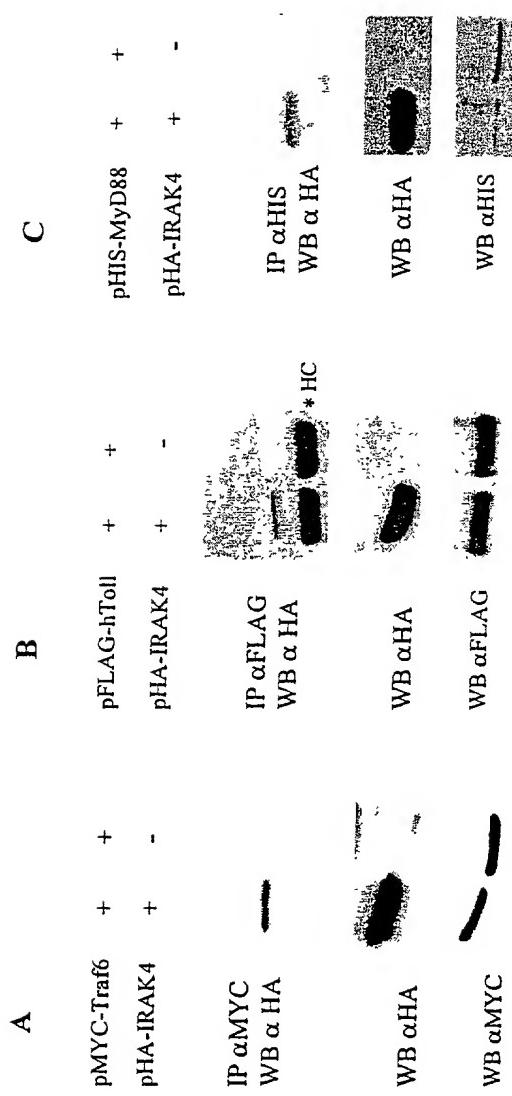


FIGURE 7

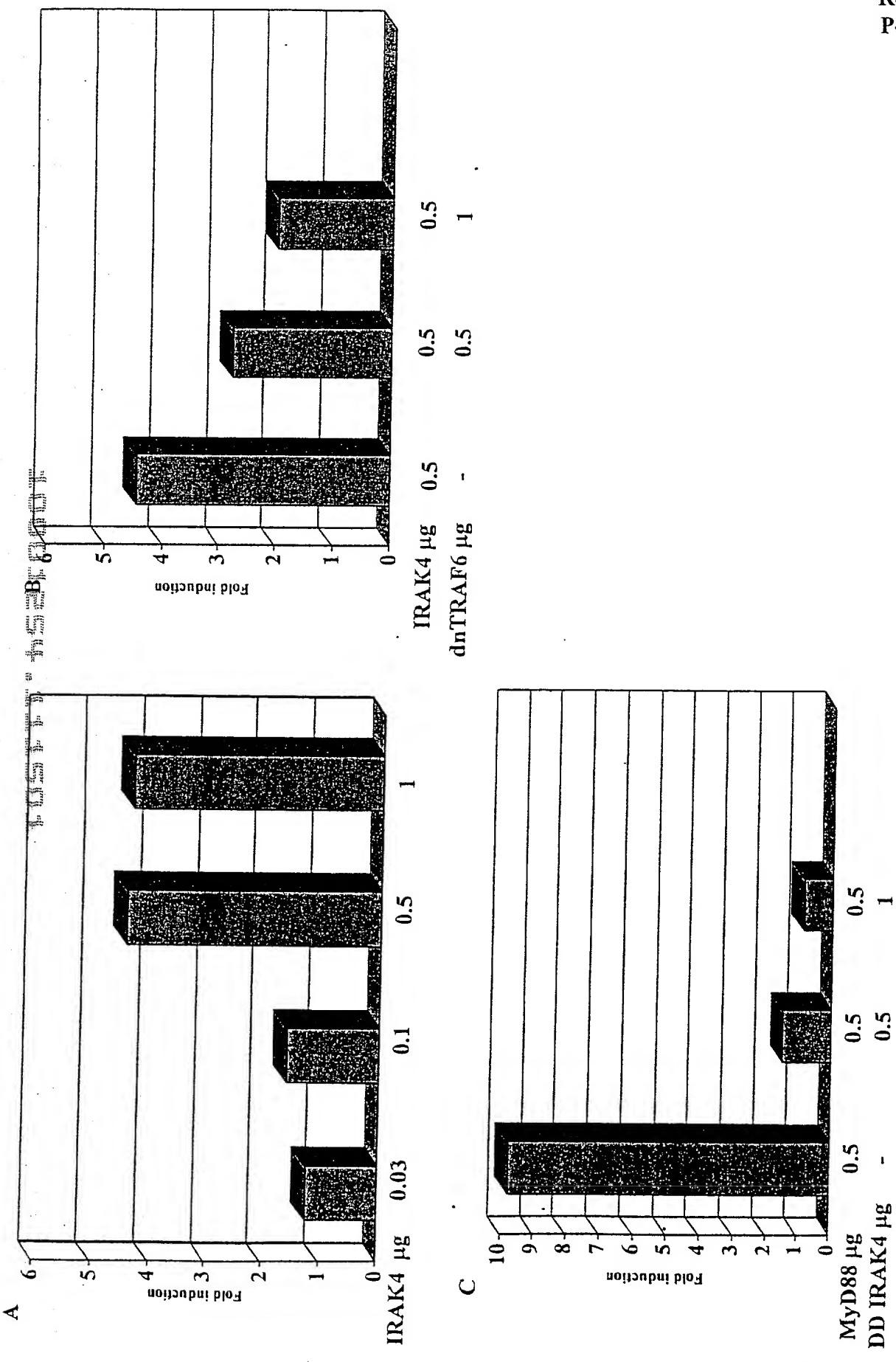


FIGURE 8

DED_4
4758144_hDEDD
6755835_mDEDD
7673638_fDedd1
PDB_FADD_1A1Z

[REDACTED] PG
[REDACTED] PG
[REDACTED] PG
[REDACTED] PG
[REDACTED] PG

DED_4
4758144_hDEDD
6755835_mDEDD
7673638_fDedd1
PDB_FADD_1A1Z

AAGG-----[REDACTED] A-----[REDACTED] GQCDESNE [REDACTED]
HE[G-----[REDACTED] NG-----[REDACTED] GQCDESNE [REDACTED]
HE[G-----[REDACTED] NG-----[REDACTED] GQCDESNE [REDACTED]
PEGSPQANTPCP [REDACTED] SQ [REDACTED] EGCSLAS [REDACTED] P [REDACTED] SGVETEELLE [REDACTED] GYISDANL [REDACTED]
[REDACTED] [REDACTED] MOSGEDDPSMTEOND [REDACTED] PPGHTEETH

DED_4
4758144_hDEDD
6755835_mDEDD
7673638_fDedd1
PDB_FADD_1A1Z

[REDACTED]
[REDACTED]
[REDACTED]
[REDACTED]
[REDACTED]

FIGURE 9

a tgaacaaacc cataacacca tcaacatatg tgcgctgcct caatgttgaa ctaatttagga agctgtcaga
tttattgtat cctcaagaag gatggaaagaa gttagctgtat gctattaaaa aaccatctgg
tgatgataga tacaatcagt ttccacataag gagattgaa gcattacttc aaactggaaa
aagtcccact tctgaattac tgtttgactg gggcaccaca aattgcacag ttggatgt
tgtgatctt ttgatccaaa atgaattttt tgctcctgcg agtctttgc tcccgatgc
tgttcccaaactgctaata cactaccttc taaagaagct ataaacagttc agcaaaaaaca
gatgcctttc tggacaaag acaggacatt gatgacacct gtgcagaatc ttgaacaaag
ctatatgcca cctgactcct caagtcaga aaataaaagt ttagaagttt gtgatacacg
ttttcacagt ttttcatttt atgaattgaa gaatgtcaca aataactttg atgaacgacc
catttctgtt ggtggtaata aaatggaga gggaggattt ggagttgtat ataaaggcta
cgtaaataaac acaactgtgg cagtgaaagaa gcttgcagca atggttgaca ttactactga
agaactgaaa cagcagttt atcaagaaat aaaagtatg gcaaaagtgtc aacatgaaaa
cttagtagaa ctacttggtt tctcaagtga tggagatgac ctctgcttag tatatgttta
catgcctaat ggttcattgc tagacagact ctctgcttg gatggactc caccactttc
ttggcacatg agatgcaaga ttgctcaggg tgcagctaat ggcataatt ttctacatga
aaatcatcat attcatagag atataaaag tgcaaataatc ttactggatg aagctttac
tgctaaaata tctgactttg gccttgcacg ggcttctgag aagtttgc agacagtc
gactagcaga attgtggaa caacagctt tatggcacca gaagcttgc gtggagaaat
aacacccaaa tctgatattt acagctttgg tggatgttta ctagaaataa taactggact
tccagctgtg gatgaacacc gtgaacctca gttattgcta gatattaaag aagaaattga
agatgaagaa aagacaattt aagatttat tggataaaag atgaatgtat ctgattccac
ttcagttgaa gctatgtact ctgttgctag tcaatgtctg catgaaaaga aaaataagag
accagacatt aagaaggatc aacagctgct gcaagagatg acagcttctt aa

FIGURE 10A

MNKPITPSTYVRCLNVGLIRKLSDFIDPQEGWKKLAVAIIKKPSG
DDRYNQFHIRRFEALLQTGKSPTSELLFDWGTNTVGDLVDLLIQNEFFAPASLLP
DAVPKTANTLPSKEAITVQQKQMPFCDKDRTLMTPVQNLEQSYPDPSSSPENKSLEV
SDTRFHSFSFYELKNVTNNFDERPISVGGNKMGEGGFGVVKGYVNNTTVAVKKLAAM
VDITTEELKQQFDQEIKVMAKCQHENLVELLGFSSEGDDLCLVYVYMPNGSLLDRLSC
LDGTPPLSWHMRCKIAQGAANGINFLHENHHIHRDIKSANILLDEAFTAKISDFGLAR
ASEKFAQTVMTSRIVGTTAYMAPEALRGEITPKSDIYSFGVVLLEIITGLPAVDEHRE
PQLLLDIKEEIEDEEKTIEDYIDKKMNDADSTSVEAMYSVASQCLHEKKNRPDIKKV
QQLLQEMTAS

FIGURE 10B

Death domains:

C. pneumoniae

DLWRQFALSLGVSEELANHEFSQAAQDMVATFRRLCDMPQLAVGLGALYTYEIQIPQVCVEKIR

C. muridarum

DLWKQFVFALGVSEELEAHEPSEAAKAKVATFMRWCTGDSLAAAGVAALYSYESQIPCVAKEKIR

C. psittaci

DLWKNFAYALGVTEEELENHVPAAAQKKVDTFLRWCTGDSLSSAGVAALYTYESQIPTVAETKIS

FIGURE 11A

GI:7190927

ORGANISM Chlamydia muridarum
ORIGIN

1 mesrkgihev smnfldqlida iiqnkhmleh pfymkwske ltkeqlqaya
kdyylhikaf

61 pkylsaihsr cddlearlkll ldnlmdeeng ypnhidlwkq fvfalgvsse
eleahepsea

121 akakvatfmr wctgdsllaag vaalysyesq ipcvakekir gleyfgfsn
pedayayftek

181 eeadvrhare ekaliemlsr ddsdkvleas revtqslygf ldsfilepatc
chchka

GI:7468151

ORGANISM Chlamydophila pneumoniae [Chlamydia pneumoniae]
ORIGIN

1 mtswielldk qiedqhmlkh efyrqwsegk lekqqqlqaya kdyylhikaf
pcylsalhar

61 cddlqirrqi lenlmdeeag npnhidlwrq falslgvsee elanhefsqa
aqdmvatfrr

121 lcdmpqlavg lgalytyeiq ipqvcvekir glkeyfgvsr rgyayftvhq
eadikhasee

181 kemlqtlvgr enpdavlqgs qevldtlwnf lssfinstep csck

FIGURE 11B

6405 ttaagc tttgtgacaa
6421 tgacaacatg ttgcaggctc taaaaatgaa tccaaaaaagc cgtataaaga ttgtgttaact
6481 tctcgcaag cttctaaaac ttgtcgcta tcatctctag acaaacatctc aattaaggcc
6541 ttttcttccc tagcatgacg cacatcagct tcttcatgct ccgtgaaata agcataatct
6601 tcaggattag aaaagccaaa gtactcaatc aatccacgaa tttttcttt agctacgcaa
6661 ggaatttgac tttcataaga atacaaagcc gctactcctg ctgctaaaga atcccctgtg
6721 caccaccgca taaatgtcgc aaccttagct ttagctgctt cactgggttc atgagcttct
6781 agctcttctg aagacactcc aagagccaaac acaaattgtt tccataaaatc aatatgatta
6841 ggataaccat tctcttcatc cattaagtta tctaataata acttgcgggc ttctaaatca
6901 tcacaacggc tatgaatagc agataaatat tttggaaaag ctttgatatg caaatagtaa
6961 tctttgcgt atgcctgtaa ttgttcttt gtcagctctc cttttgacca cttcatgtaa
7021 aaagggtgtt ctaacatatg tttgtttga ataattgcat cttagctgatc taaaaaattc
7081 atgctcacct cttttattcc ttttcttgat tccac 7115

FIGURE 11C

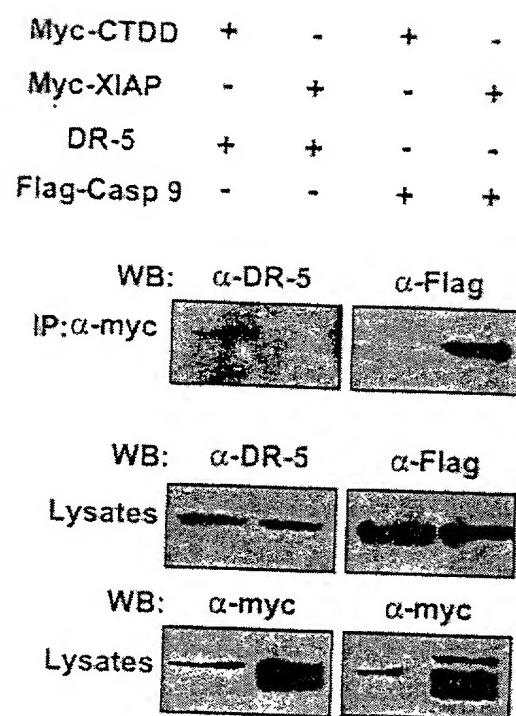


FIGURE 12

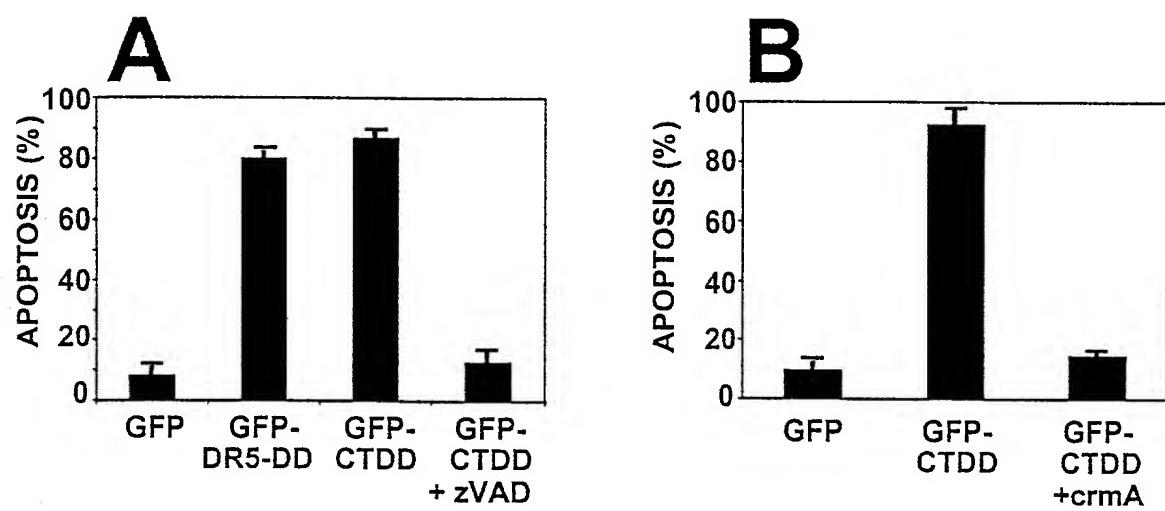


FIGURE 13

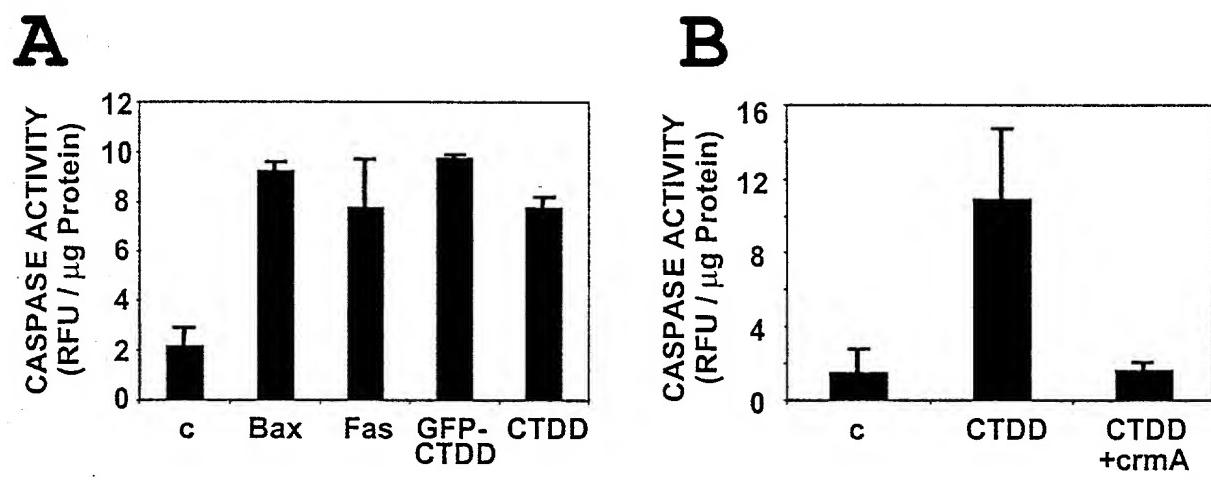


FIGURE 14

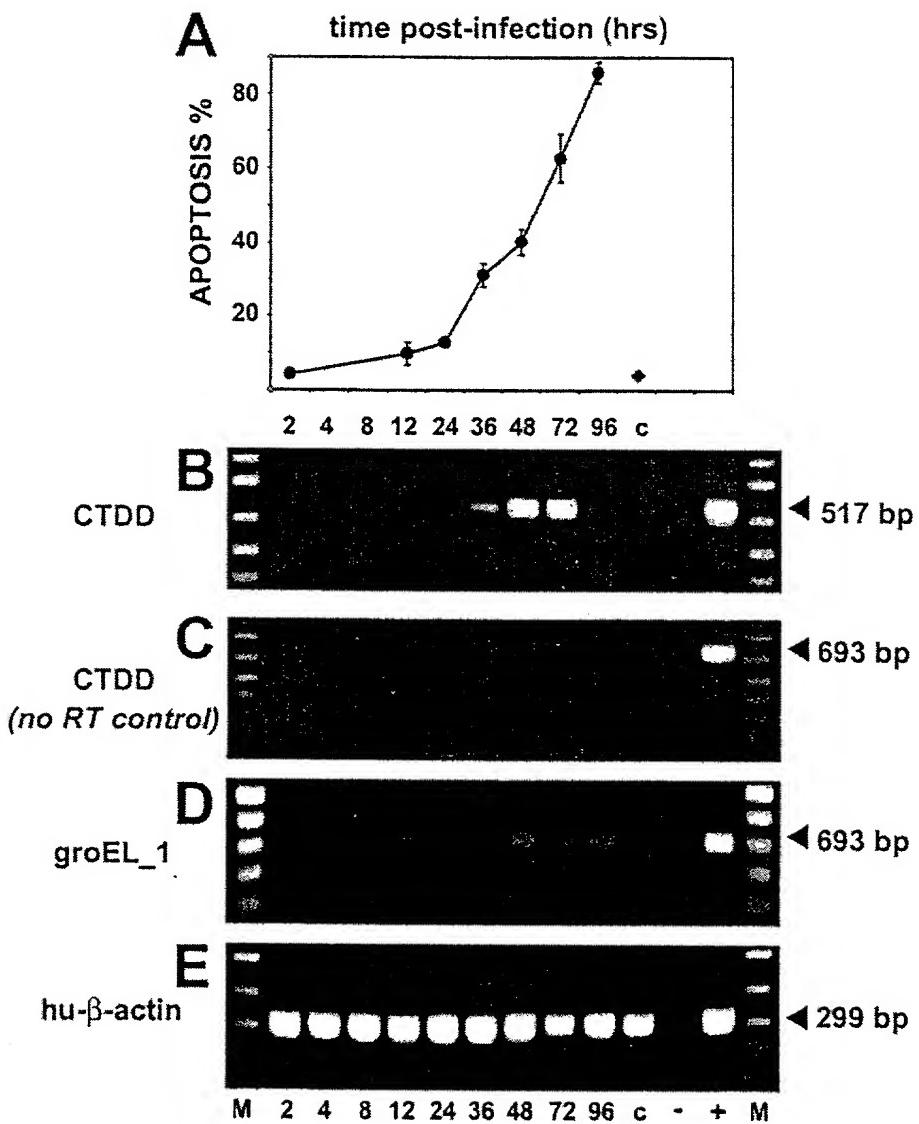


FIGURE 15